

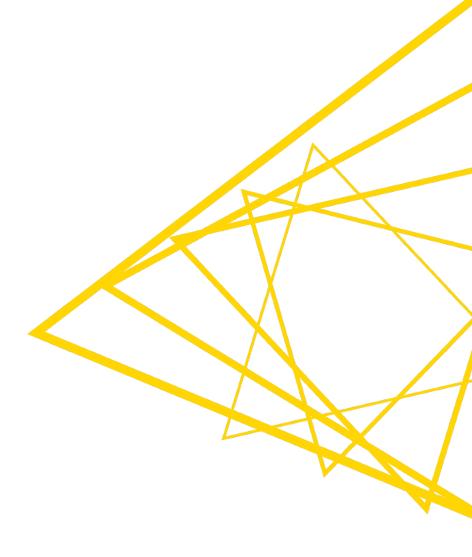
Artificial Intelligence

Visual Programming for Data Science

Working with Data at the right Level of Abstraction

Prof. Dr. Michael Berthold ISACA Fokus Event, Nov 10, 2022

Data Science Personas



Data Science Researchers

Researchers focus on Data Science algorithms

- often have CS/NumMath/Stats background
- use Python, Javascript, R, SQL, C, Java, or XYZ
- tend to not worry about others.

```
## On my machine, it takes 39 seconds with 1 worker and 10 seconds with 4.
                                                                                                                                                                                                  ## 29.055u 0.102s 0:28.68 101.6% 0+0k 0+3io 0pf+0w
                                                                                                                                                                                                  #max workers=1
                                                                                                                                                                                                  ## With 4 threads it takes 11 seconds.
#include <iostream>
                                                                                                                                                                                                  ## 34.933u 0.188s 0:10.89 322.4% 0+0k 125+1io 0pf+0w
                                                                                                                                                                                                  max workers-4
#include <segan/file.h>
#include <segan/seguence.h>
                                                                                                                                                                                                   # (The "u"ser time includes time spend in the children processes.
                                                                                                                                class TICWritingConsumer : public MSDataWritingConsumer
                                                                                                                                                                                                  # The wall-clock time is 28.68 and 10.89 seconds, respectively.)
using namespace segan;
                                                                                                                                                           WritingConsumer allows to change the . # This function is called in the subprocess.
                                                                                library(foreach)
                                                                                                                                                           isk (to "filename") using the processS # The parameters (molecule and number of conformers) are passed via a Python
                                                                                                                                                                                                  def generateconformations(m, n):
                                                                                length divisor<-6
int computeLocalScore(String<char> subText, String<char> pattern)
                                                                                                                                                                                                       - Chem. AddHs (m)
                                                                                iterations<-5000
                                                                                                                                                                                                     ids=AllChem.EmbedMultipleConfs(m, numConfs=n)
    int localScore = 0;
                                                                                predictions <- foreach (m=1:iterations,.combine=cbind)
                                                                                                                                                                                                        AllChem.UFFOptimizeMolecule(m, confId=id)
    for (unsigned i = 0; i < length(pattern); ++i)</pre>
                                                                                                                                                                                                     # EmbedMultipleConfs returns a Boost-wrapped type which
                                                                                                                                                            et TIC to zero
         if (subText[i] == pattern[i])
                                                                                                                                                                                                     # cannot be pickled. Convert it to a Python list, which can.
                                                                                                                                                           g filename) : MSDataWritingConsumer(fi
                                                                                                                                                                                                     return m list(ids)
              ++localScore;
                                                                                training positions <- sample(nrow(training),
                                                                                                                                                                                                  smi_input_file, sdf_output_file = sys.argv[1:3]
                                                                                size=floor((nrow(training)/length divisor)))
                                                                                                                                                           step for spectra before they are writ-
    return localScore;
                                                                                train pos<-1:nrow(training) %in% training positions
                                                                                                                                                                                                  n = int(sys.argv[3])
                                                                                                                                                           ataWritingConsumer::SpectrumType & s)
                                                                                lm fit<-lm(y~x1+x2+x3,data=training[train pos,])</pre>
                                                                                                                                                                                                  writer = Chem.SDWriter(sdf_output_file)
                                                                                                                                                           .size(); i++) { TIC += s[i].getIntensi
String<int> computeScore(String<char> text, String<char> pattern)
                                                                                predict(lm fit,newdata=testing)
                                                                                                                                                                                                  suppl = Chem.SmilesMolSupplier(smi_input_file, titleLine=False)
                                                                                                                                                                                                  with futures.ProcessPoolExecutor(max workers-max workers) as executor:
    String<int> score;
                                                                                                                                                           .(MSDataWritingConsumer::ChromatogramTy
                                                                                                                                                                                                     # Submit a set of asynchronous jobs
                                                                                predictions <- rowMeans (predictions)
    resize(score, length(text) - length(pattern) + 1, 0);
                                                                                error <- sqrt ((sum ((testing $y-
                                                                                                                                                                                                     for mol in suppl
                                                                                                                                                                                                        if mol:
    for (unsigned i = 0; i < length(text) - length(pattern) + 1;</pre>
                                                                                                                                                           har** arqv)
                                                                               predictions)^2))/nrow(testing))
                                                                                                                                                                                                            job = executor.submit(generateconformations, mol, n)
         score[i] = computeLocalScore(infix(text, i, i + length(pattern,,,
                                                                                                                                                                                                            jobs.append(job)
                                                                                                                                     (argc < 2) return 1
                                                                                                                                  // the path to the data should be given on the command line
                                                                                                                                                                                                     widgets = ["Generating conformations; ", progressbar.Percentage(), " ",
                                                                                                                                  String tutorial_data_path(argv[1]);
                                                                                                                                                                                                              progressbar.ETA(), " ", progressbar.Bar()]
    return score;
                                                                                                                                                                                                     pbar = progressbar.ProgressBar(widgets=widgets, maxval=len(jobs))
                                                                                                                                  // Create the consumer, set output file name, transform
                                                                                                                                                                                                     for job in pbar(futures.as_completed(jobs)):
                                                                                                                                  TICWritingConsumer * consumer = new TICWritingConsumer("Tutori
                                                                                                                                                                                                         mol.ids=iob.result()
int main()
                                                                                                                                                                                                         for id in ids:
                                                                                                                                  MzMLFile().transform(tutorial_data_path + "/data/Tutorial_FileI(
                                                                                                                                                                                                            writer.write(mol, confId=id)
                                                                                                                                  std::cout << "There are " << consumer->nr.spectra << " spectra
    String<char> text = "This is an awesome tutorial to get to know SeqAn!";
                                                                                                                                  std::cout << "The total ion current is " << consumer->TIC << std::endl;
    String<char> pattern = "tutorial";
                                                                                                                                  delete consumer;
    String<int> score = computeScore(text, pattern);
                                                                                                                                  return 0;
                                                                                                                                } //end of main
    for (unsigned i = 0; i < length(score); ++i)
```



""" contribution from Andrew Dalke """

DownLoad this from http://pypi.python.org/pypi/futures

Download this from http://pypi.python.org/pypi/progressbar

from rdkit.Chem import AllChem

from concurrent import futures

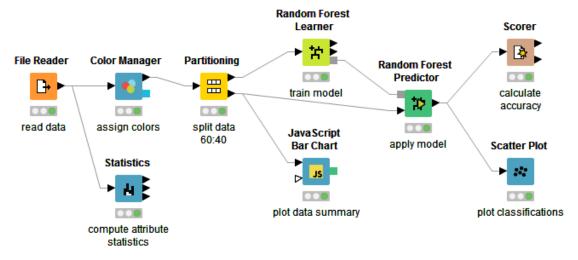
import sys

import progressbar

Data Scientists

Data Scientists

- have (also) a specific domain background
- often can program in Python, R, SQL, C, Java, or ...
- ...but really focus on the data flow
- want to use and try out other stuff

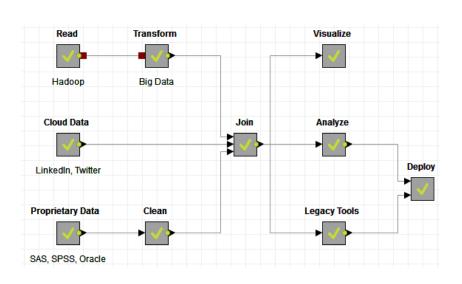


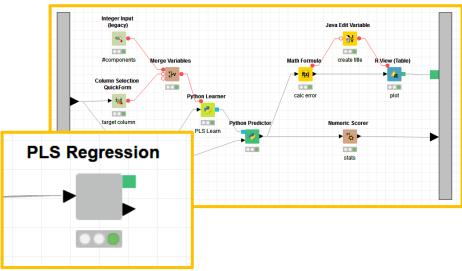


Casual Users ("Citizen Data Scientists")

The occasional data person

- wants to start from something existing that solves a similar problem
- often comes from an Excel, BI, or other data world
- needs solid documentation and proper abstraction







Data Science Consumers

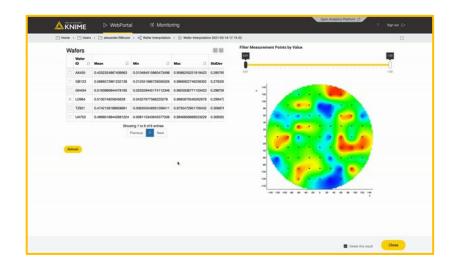
Data Science Consumers use

their favorite frontend
 (= Data Science needs to embedded via an API)

or

Interactive Analytics (= Data Science Applications)







Data Science Personas

library(foreach)
lampa(s)(s)one-d
lampa(

Coders: Programs & Scripts

Data Science
Algorithm Inventors



Data Scientists: Visual Programming

Data Science Creators



Citizen Data Scientists: Components & Blueprints

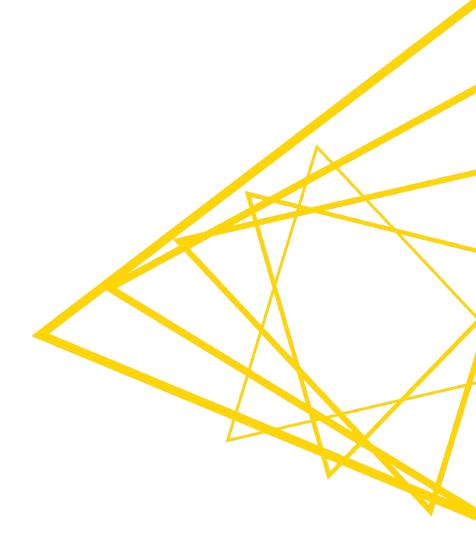


Data Science Consumers: Applications & Services

Data Science Consumers



No Code, Low Code, Visual Programming?



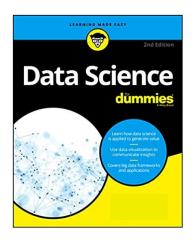
No Code

No Code for Simple Problems:

No Code

(Code-Based) Programming

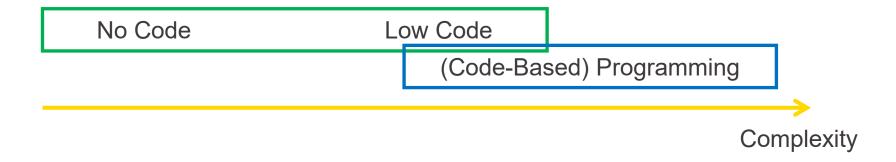
Complexity



Visual Environment allows quick creation of solutions for standard problems.

Low Code

Low Code for Standard Problems:



Visual Environment allows 80% creation of solutions.

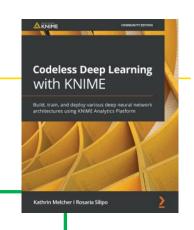
...for the rest (& details): reach out to code.

(often: Visual UI on top of **one** programming language.)

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Visual Programming

Visual Programming for all types of Problems:



de EXCEI para KNIME ©
Autors: Kathrin Melcher
Tabulati Maria Basse di basa Cost Autors feesald: Catalor Peesa Bassa. Cost Autors feesa Bassa. Cost Autors feesa

Visual Programming

(Code-Based) Programming

Complexity

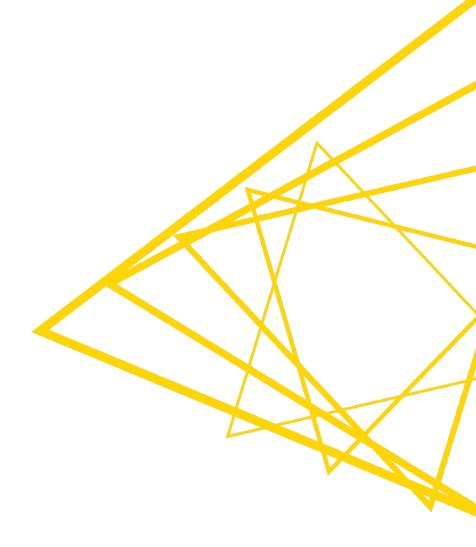
Visual Environment allows complete creation of solutions. (from simple to complex)

If desired: you can also include code (of various languages!).

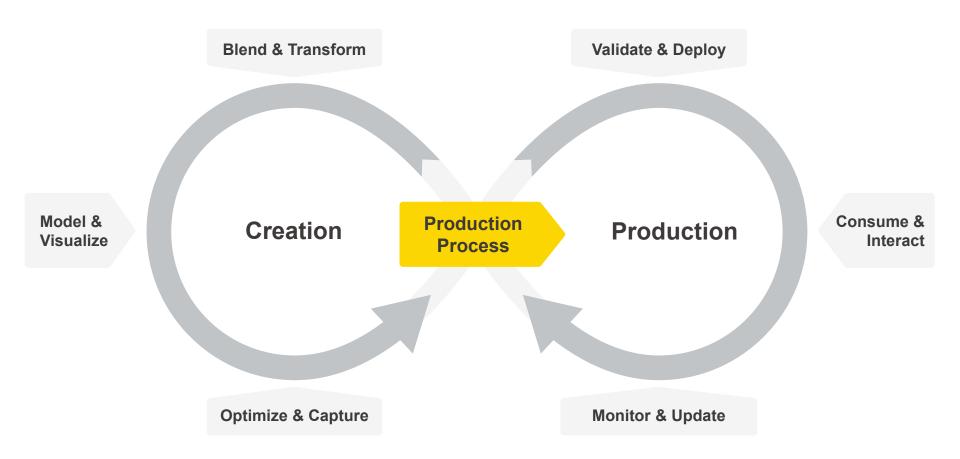
(Visual UI models what's running under the hood.)

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Visual Programming for Data Science



The Data Science Life Cycle



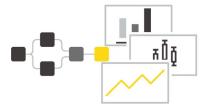
Data Science Activities



Data Wrangling (Virtual Warehouses)



Descriptive Analytics (Automating Repetition)



Diagnostic Analytics (Creating Insights)

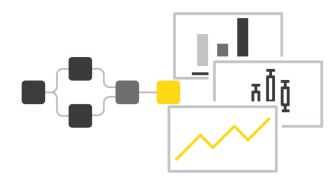


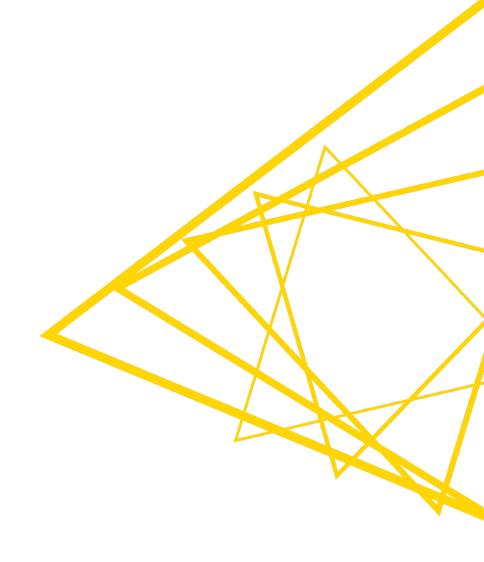
Advanced Analytics (AI/ML...) (Predictive, Prescriptive)



Productionize (Deploy, Manage, Govern)

No Code for Analytics

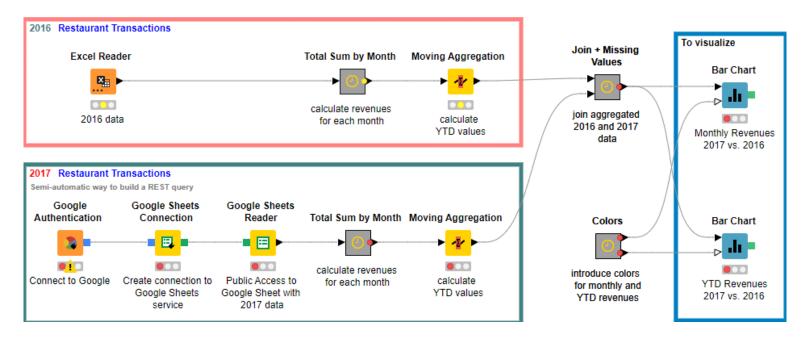




No Code for Analytics

Benefits:

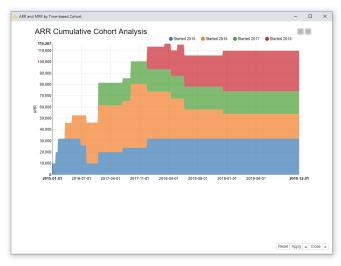
- (Reliable) Automation & Reproducibility
- Documentation



Creating Data Insights: Abstraction helps

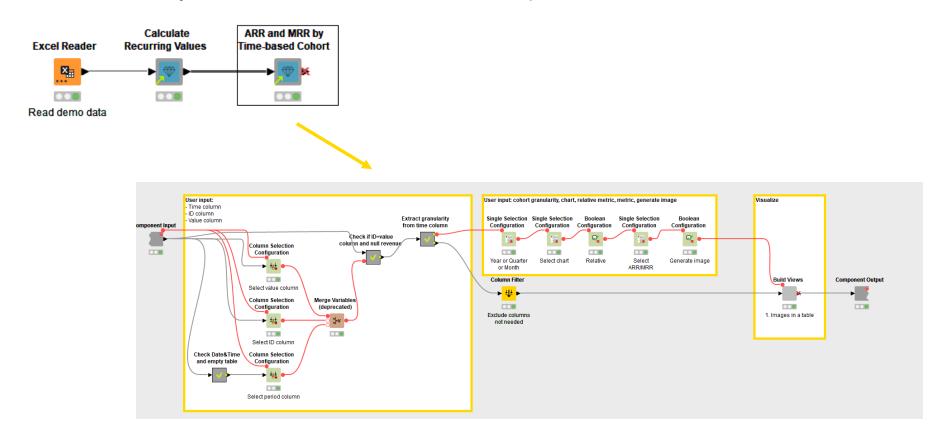
ARR Cohort Analysis for Excel Users





Creating Data Insights: Abstraction helps

ARR Cohort Analysis for Excel Users – No Code Encapsulated:



Abstraction in KNIME: Components



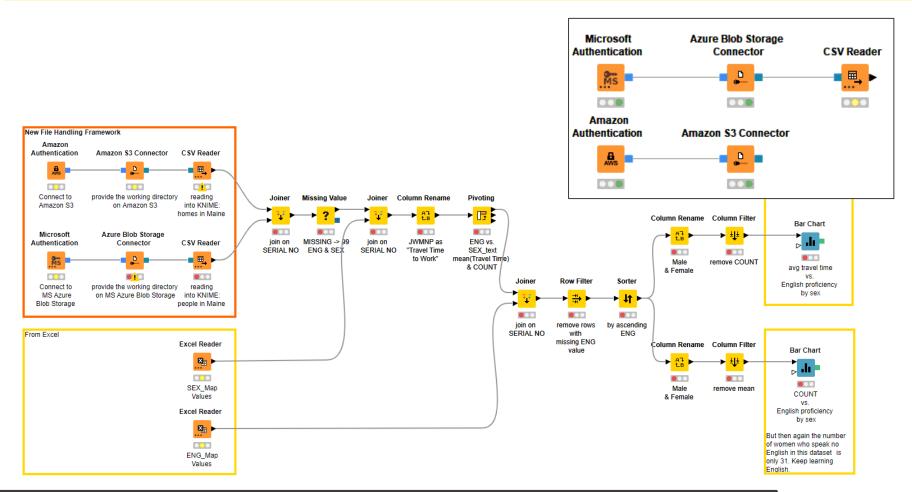


Low Code for Data Wrangling



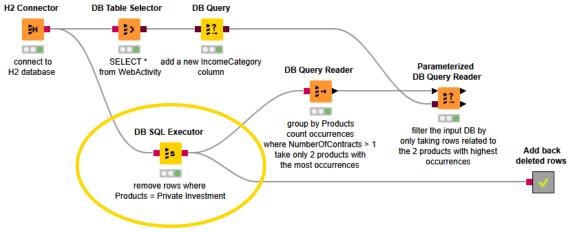


Flexible Data Wrangling



KI

Flexible Data Wrangling

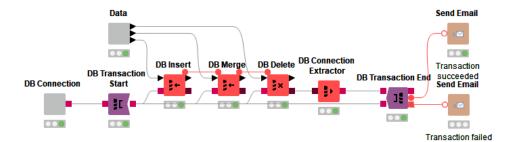


- ▼ .º Connectors
 - Amazon S3 Connector
 - Azure Blob Storage Connector
 - Azure Data Lake Storage Gen2 Connector
 - Databricks File System Connector
 - FTP Connector
 - Google Cloud Storage Connector
 - Google Drive Connector
 - HDFS Connector
 - HDFS Connector (KNOX)
 - L HTTP(S) Connector
 - E KNIME Server Connector
 - KNIME Workflow Data Area Connector
 - Local File System Connector
 - Ms Microsoft Authentication
 - SMB Connector
 - SSH Connector
 - SharePoint Online Connector

- ▼ **3** DB
 - ▼ Connection
 - A Amazon Athena Connector
 - R Amazon Redshift Connector
 - DB Connection Closer
 - DB Connection Extractor
 - DB Connector
 - ₽ H2 Connector

 - : Impala Connector

 - A Microsoft Access Connector
 - s Microsoft SQL Server Connector
 - MySQL Connector
 - **3**o Oracle Connector
 - PostgreSQL Connector
 - SQLite Connector
 - s Snowflake Connector
 - ▼ Vertica Connector



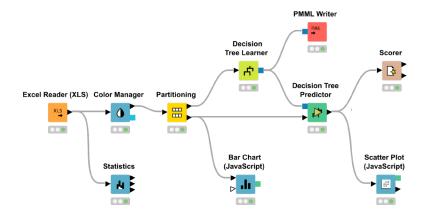
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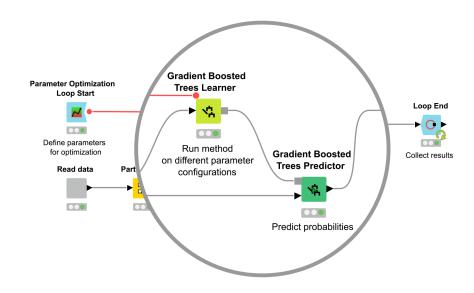
Low Code for Data Science





No Code for Data Science



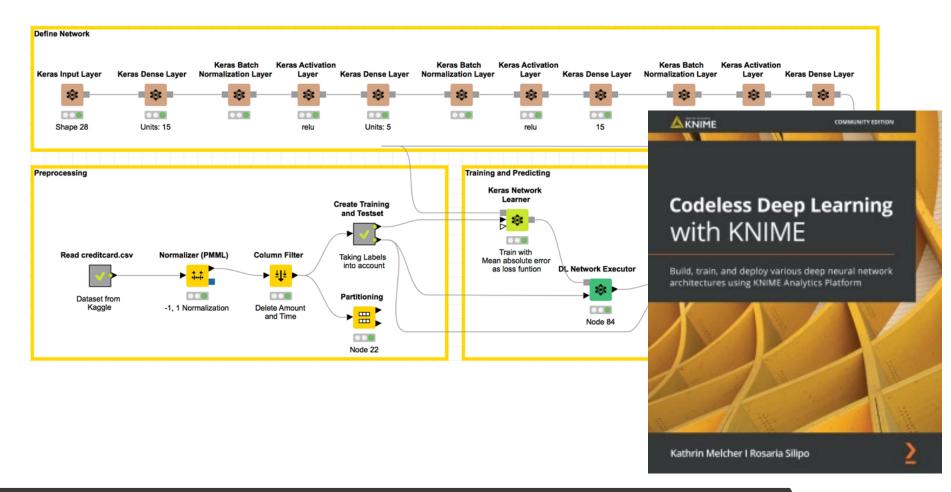




No Code for Interactive Visualizations



Codeless Deep Learning

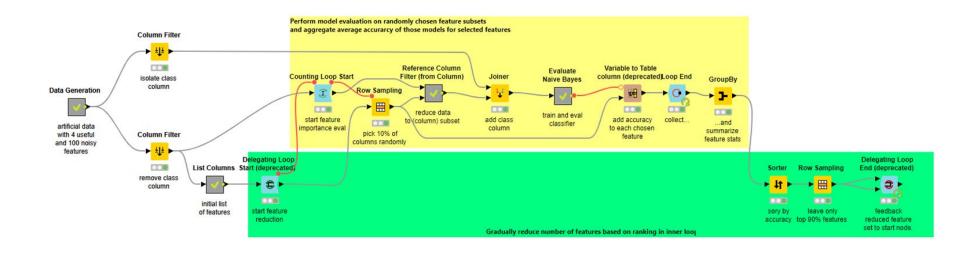


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Data Science *is* complex: Visual Programming

Visual Data Science Algorithms: Recursive Feature Elimination



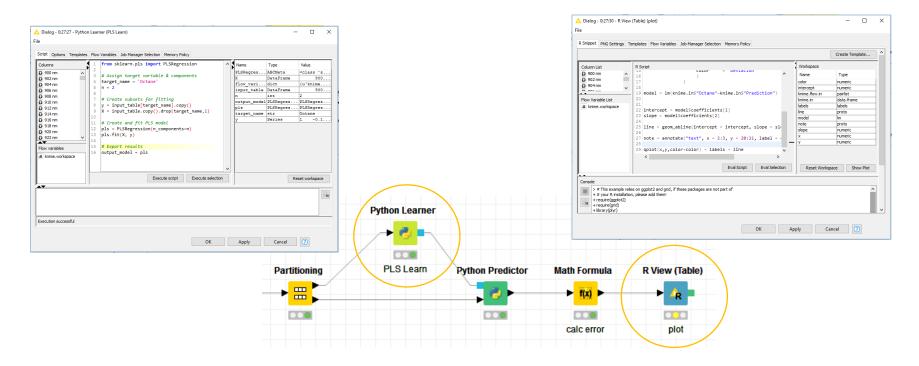
"Computer programming is an art, because it applies accumulated knowledge to the world, because it requires skill and ingenuity, and especially because it produces objects of beauty."

Donald Knuth 1974



Low Code for Data Science

Leveraging Code in Visual Programming:



...works with (and from...) Jupyter Notebooks as well.

Data Science *is* complex: More Topics

Data Science "Stuff":

Data Blending:

- Feature Selection
- Feature Engineering
- Anonymization
- · ...

Modeling

- Model Selection
- Parameter Optimization
- Ensemble Creation
- Active Learning
- Transfer Learning
- •

Management Requirements:

Compliance

- Best Practices
- Required Standards
- Bias Detection and Removal
- Data Privacy / GDPR
- Explainable "AI"
- ...

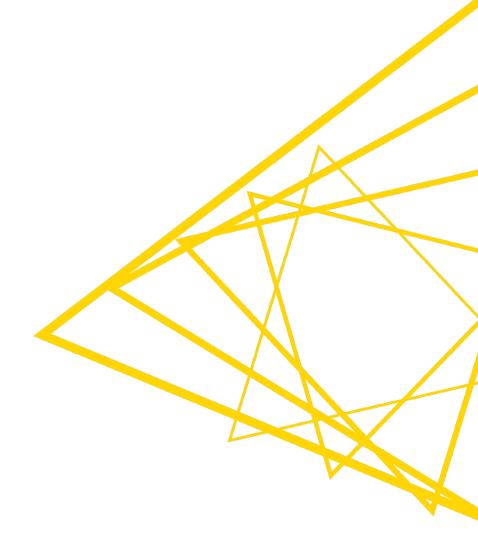
Governance

- Integration / Dependencies
- Traceable / Lineage
- Documented
- Reproducibility
- ...

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Data Science Teams





The Data Science Life Cycle

Blend & Transform



Model & Visualize



Deploy & Manage



Consume & Interact



Techniques:

- Joins
- Pivot
- Normalization
- PCA
- ...

- Clustering
- Regression
- Deep Learning
- Visualizations
- ..

- Model Monitoring
- Updating
- Validation
- ...

- Active Learning
- Interactive Applications
- Services
-

Technologies and Languages:

- • R, Python
- Data Science Teams shouldn't really have to worry about all of this...
- Graf

- Diotly, R-Shiny JavaScri<mark>ot</mark> Vue.js
- Grafana RES

...

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Democratizing Data Science

What's the Essence of the Job:

Create (complex) Data Science processes, often together with other Experts

What's usually not part of the Job:

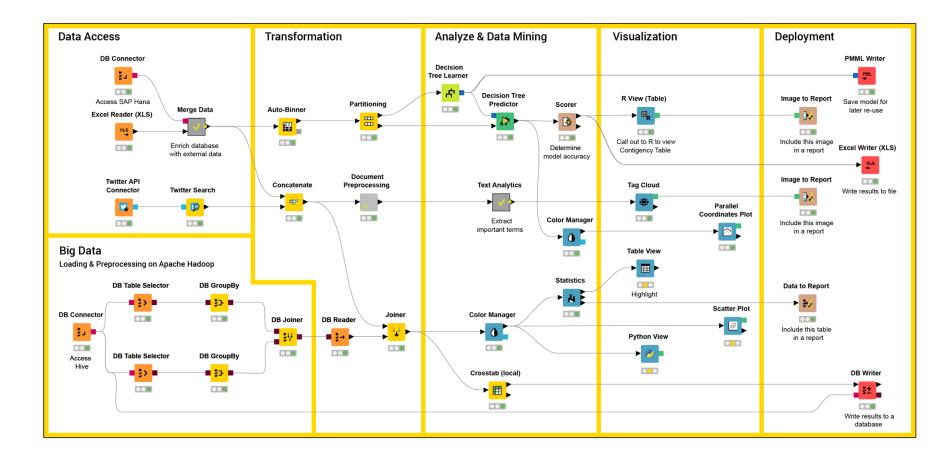
Inventing, Writing, Optimizing new Algorithms

What shouldn't be part of the Job:

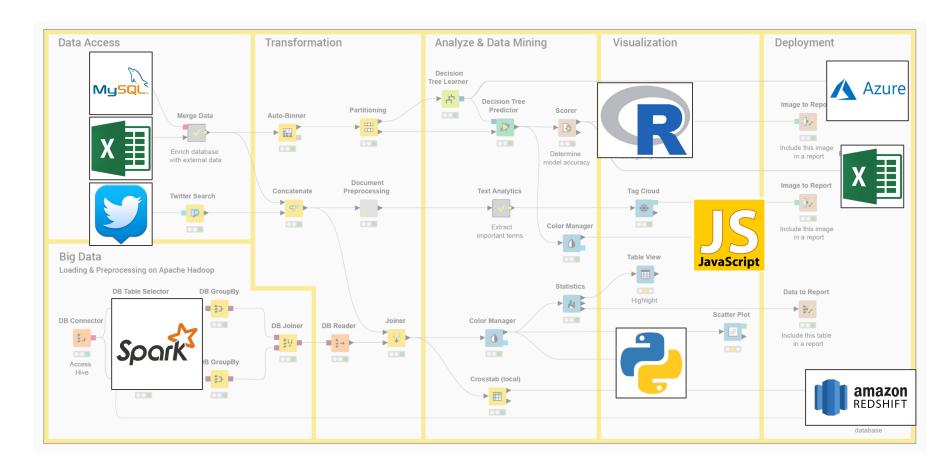
- Caring about the details of the underlying technology
- Worrying about interfaces of different tools
- Worrying about library versions and (backwards) compatibility
- Manually doing repetitive tasks

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Visual Programming for Data Science

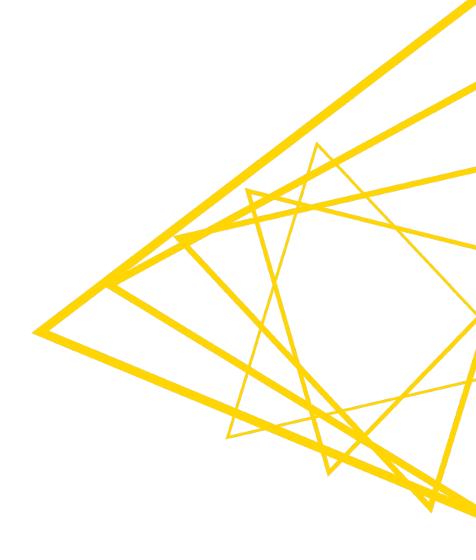


Technologies & Languages under the Hood



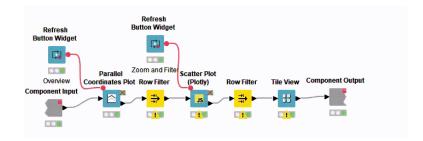
Low Code for Deployment



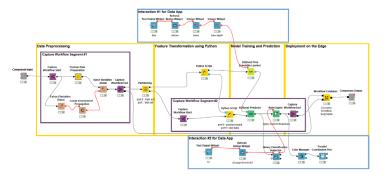


No Code Deployment of Data Apps

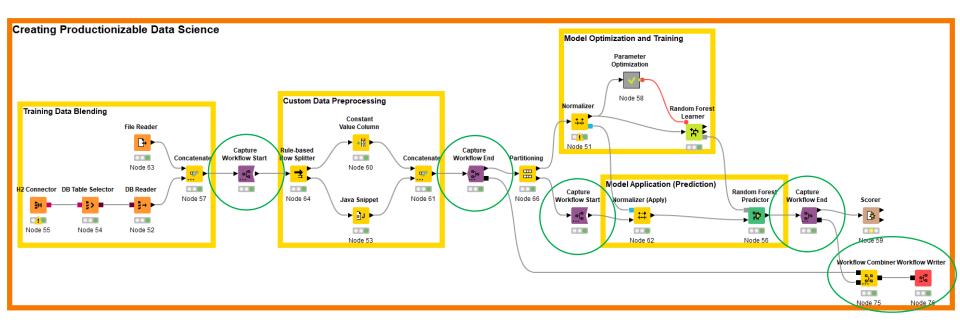






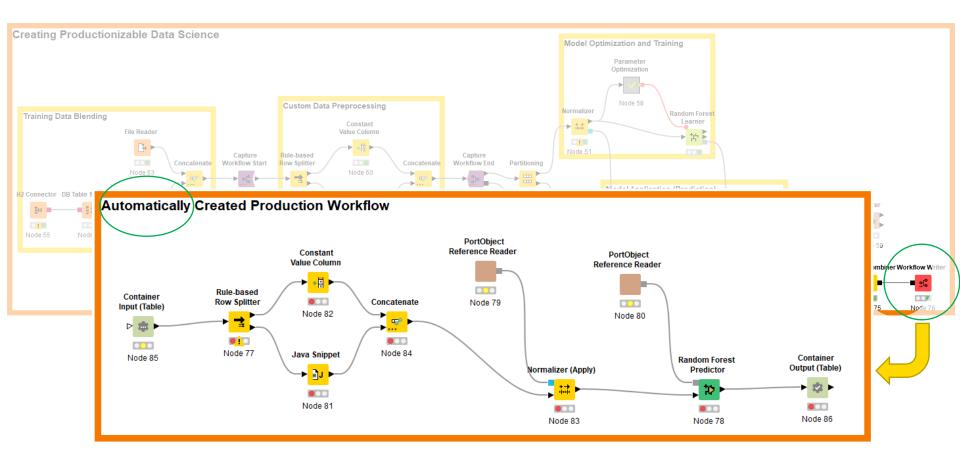


Deploying Data Science as Service





Deploying Data Science as Service





Visual Programming for all Aspects of Data Science



Data Wrangling (Virtual Warehouses)



Descriptive Analytics (Automating Repetition)



Diagnostic Analytics (Creating Insights)



Advanced Analytics (AI/ML...) (Predictive, Prescriptive)



Productionize (Deploy, Manage, Govern)



Visual Programming for Data Science makes Sense!

- Data Science is a Team Sport:
 - requires a broad spectra of skills often done in collaboration
 - requires understanding and access to the inner wheels of an algorithm (but not to the algorithm itself)
 - No single language or tool has all the answers it's about tool blending as well!
- No-Code, Low-Code, Visual Programming (if done right):
 - automates repetitive tasks
 - provides the appropriate level of abstraction to allow focus on modeling a data process
 - allows access to all relevant nuts and bolts
 - allows to abstract / encapsulate sophistication for others to safely reuse
 - removes tool interaction/interface complexity
 - provides transparency & governance
 - (and it's not just a UI slapped on top of a language)
 - ...allows everybody to gradually increase their Data Science Proficiency (in the same Environment).



Data Science Upskilling

Individual Sophistication

NIL & Al

Stats and Models

Data Exploration

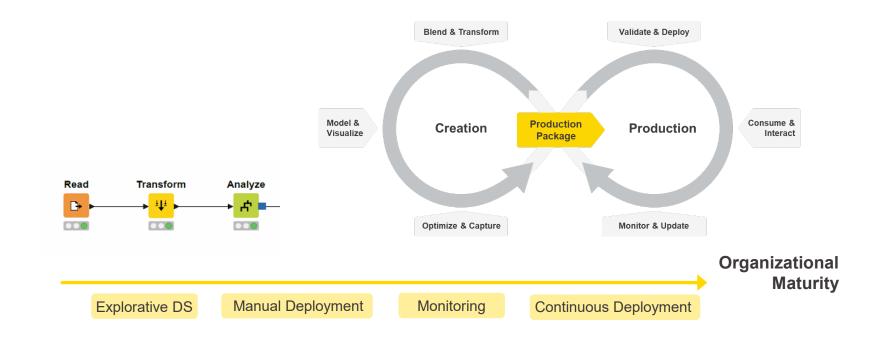
Data Exploration

Automated ETL

Apreadsheets & Bl

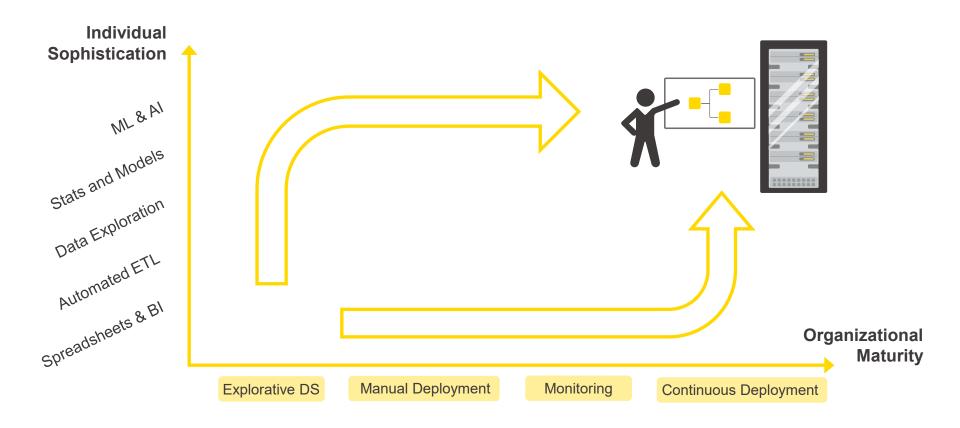


Organizational Upskilling



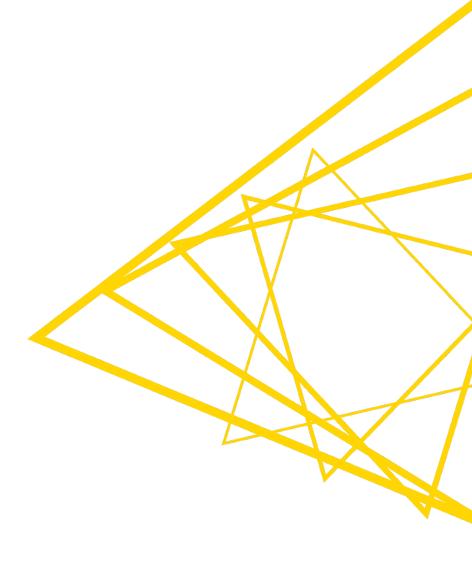
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Upskilling Individuals and Organizations





Want to get started?



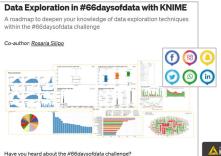
Getting Started?



Beginners corner



https://www.knime.com/learning



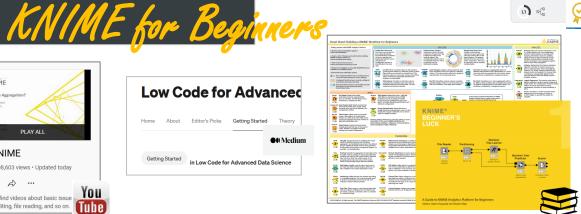














[L1-DS] KNIME Analytics Platform for

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O Course Length 8 hours

★ Rating ★★★★★ 1079 Reviews

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